CHARACTERIZATION OF THE XRCC2 HUMAN DNA REPAIR PROTEIN; N. Liu, M.P. Thelen*, J.E. Lamerdin*, W. Fan*, M. Hwang*, J.D. Tucker*, and L.H. Thompson, Lawrence Livermore National Laboratory, Livermore, CA 94551

XRCC2 is a human repair protein that corrects the hypersensitivity of hamster mutant irs1 to ionizing radiation, cross-linking agents, and other DNA damage agents. The XRCC2 cDNA, encoding at least 280 amino acids, was isolated using functional complementation. The XRCC2 genomic DNA, obtained by screening a P1 library with XRCC2 cDNA probe (GenomeSystems, Inc.), also complemented irs1 but was found to be truncated at the 5' end and missing ~80 bp in the cDNA. The XRCC2 cDNA and genomic DNA transfor-mants showed partial correction in terms of cell survival for MMC, but showed nearly full correction for cisplatin as examined by a differential cytotoxicity assay. With ionizing radiation, the genomic transformant showed wild-type sensitivity while the cDNA trans-formants surprisingly showed no correction. Chromosomal aberrations showed intermediate correction in both plasmid and genomic transformants in unirradiated cultures as well as after 100 cGy of ¹³⁷Cs γ-rays. The predicted protein sequence shows weak similarity with S. cerevisiae Rad51 and contains consensus ATP binding domains. The XRCC2 cDNA was cloned into expression vectors pET32 and pET26 (Novagen) and overexpressed in E. coli. His-tagged thioredoxin(trx)-XRCC2 fusion protein was obtained in the soluble fraction and purified to near homogeneity using a Ni-NTA agarose column (Oiagen). Insoluble pelB-XRCC2 fusion protein with C-terminal His-tag was purified in denaturing 8 M urea, gradually dialyzed to 1 M urea, and used for producing polyclonal antibodies in mice. The antisera detected both the overexpressed pelB-XRCC2 and the trx-XRCC2 fusion proteins, and they also detected a protein of ~ 50 kDa in HeLa cell extract. Thus, the current XRCC2 cDNA is probably truncated at the 5'-end. Experiments are in progress to obtain the cDNA sequence upstream of the existing cDNA as well as the missing portion of the genomic sequence. (Work was done under the auspices of the U.S. DOE by LLNL under contract No. W-7405-ENG-48).